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RESULT 5
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80)
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Brassica oleracea
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         'karyota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; rophyta; Magnoliophyta; eudicotyledons; core eudicots; rosic
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2 Brassica
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28;
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            rosids
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S., Utterback, T.,

Koo, H. and Fraser, C.M.

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180

CATTICACGITCGCTGAAGG 199

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EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Z98448
HSZ98448 DKFZphthml
                                                                                                                                                                                                            Deutsches Krebsforschungszentrum Heidelberg
Im Neuenheimer Feld 506, D-69120 Heidelberg,
Location/Qualifiers
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Mammalia; Eutheria;
1 (bases 1 to 199)
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21; Conserv
                             Similarity 100.
20; Conservative
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                                                                                                                                                                                                                                                                  Contact:
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                                                                                                                                                                                                                                                                                                         Korn, B.,
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                                                                                                                   /organism="Homo sapiens"
/mol type="mRNA"
/db xref="taxon:9606"
/clone="xkFzphthml_1e7"
/tissue_type="thymus"
/dev_stage="adult"
                                                                                        /clone___
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100.0%; Pr
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                            Score 20; DB; Pred. No. 80; 0; Mismatches
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HONINTO_PHOS1 using BstXI linkers"
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                                                     Length 199;
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Enterobacteria phage P1
Viruses; dsDNA viruses,
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AJ011592
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Nucleotide sequence of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished
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                                  /db_xref="SPTREMBL:O80281"
/translation="MGASPLESMPNSLSAEQAVLGGLMLDNCRWDEVADRIVADDFYT
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/translation="MGASPLESMPNSLSAEQAVLGGFAYLAEMSKNTPSSANIC
ANADIVRERAVVREMISVANEIAEAGYAQDGRGSNELLDMAERRVFEIAEKQKSGSG
PKDIASILDATVSRIEELFQRPHDGVTGLDTGFTDLNKKTAGLQPSDLIIVAARPSMG
KTTFAMMLVENAAVRNDKPULVFSLEMPSHQLMRSLASLAEWDGTRIRTGQLNDEDM
ARVSGAMGILLDKQNIFIDDSSALTFTELRSARRVYKENGGLSMIMIDVLQLMRVPE
LQDNRTLEIAEISRSLKALAKELQVPVVALSQLNRSLEQRADKFVNSDLRESGAIEQ
DADLLMFLYRDEVYHPDSEMKGIAEVIIGKQRNGFIGTVRLAFNGQYSRFDNYAGADM
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                      QEDY"
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/protein_id="CAA09719.1"
/db_xref="GI:3688227"
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/strain="Plcm cl.100"
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e bacteriophage P1 ban helicase gene
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/product="Ban"
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SDSVSDGDFYRHDHRLIFRAIYKLAEANQPIDVVTLSEQLEKEGQLN
AKNIPSVANIKAYAQIIRERATLRQLIGISNEIADSAFHPEGRGANE
VAEARPKTGGPVGISDILVKTIDRIDYLFNTTEALTGVSTGFTDLDEKT
IVAGRPSMGKTTFAMNLVENAVMRTDKAVLVYSLEMPSDSIVMRMLSSLGR
SGKLDDEDWPRLTSAINLLNDKKLFIDDTAGISPSEMRARTRRLVREHGDLJ
                                                                                 /transT_table=11
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/protein_id="AAKO0230.1"
/db_xref="GI:12642368"
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